

Genetic variation and molecular authentication of selected *Aquilaria* species from natural populations in Malaysia using RAPD and SCAR marker.

ABSTRACT

Aquilaria (Thymelaeaceae) is an endangered agarwood-producing tropical tree that is endemic to the Indomalesia region. Molecular information on genetic diversity of *Aquilaria* is limited. The aims of this research were to study genetic diversity among three *Aquilaria* species, growing in natural and distant populations in Malaysia using RAPD markers and to develop SCAR markers for easy identification of *A. malaccensis*, the major agarwood producer. By analyzing 23 RAPD primers, a total of 368 bands were scored. Multi-populations Descriptive statistics revealed that 333 (90.49%) polymorphic bands were found at species level, where *A. malaccensis* had 107 (29.08%) bands, *A. hirta* had 56 (15.22%) and *Aquilaria* sp.1 had 11 (2.99%), for the percentage of polymorphic loci in a single population/species. Nei's unbiased measurement indicated moderate similarities among populations/species. Out of the 23 RAPD primers, three were found specific to *A. hirta* and one was specific to each *A. malaccensis* and *Aquilaria* sp.1. RAPD-based SCAR markers generated a total of five species-specific amplicons: three for *A. hirta*, one each for *A. malaccensis* and *Aquilaria* sp.1. SCAR markers for *A. malaccensis* were used to distinguish five other different *A. malaccensis* populations in Malaysia. SCAR markers for *A. malaccensis* tested in five other different *A. malaccensis* populations in Malaysia yielded positive and consistent results. The DNA fingerprints identified for each *Aquilaria* sp. will be useful for *Aquilaria* identification in natural population, young plantation and even at seedling and seed stages in the nursery, as it is rapid and cost-effective and does not rely on morphology.

Keyword: Agarwood; *Aquilaria*; Genetic variation; Genetic distance; Molecular markers.